SEQUENCE LISTING

FANG KOHLSTAEDT, LORI RENO, JOHN

<120> HUMANIZED ANTIBODIES

<130> 014357/027 8772

<140> 09/910,483 <141> 2001-07-19

<160> 96

<170> PatentIn Ver. 2.1

<210> 1 <211> 116

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Synthetic VH Domain peptide of Hum A

Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Asn Ile Lys Asp Thr 20 25 30

Tyr Ile His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val

Ala Arg Ile Asp Pro Ala Asn Asp Asn Thr Ile Tyr Ala Asp Ser Val

Lys Gly Arg Phe Thr Ile Ser Ser Asp Asp Ser Lys Asn Thr Ala Tyr 65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr 85 90

Thr Asp Ser Gly Tyr Trp Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val

Thr Val Ser Ser 115

<210> 2 <211> 348 <212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic VH nucleotide sequence of Hum A

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gctgacagcg tgaagggccg ttttactatt tctagcgacg actctaagaa caccgcgtac 240
cttcagatga actctctgcg tgccgaggac accgccgtct actactgcac ggactctggc 300
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      peptide of Hum A
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Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Asn Asn
Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
35 40 45
Tyr His Ala Ser Gln Ser Ile Ser Gly Val Pro Ser Arg Phe Ser Gly
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Asn Ser Trp Pro Tyr
Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg
<210> 4
<211> 324
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic VL nucleotide
      sequence of Hum A
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attacttgcc gcgccagcca atctatcagt aataatcttc actggtatca acaaaaaccg
ggtaaagčtc čgaaacťtct tatctatcac gcctctcaga gcaťťagcgg cgttccgagc 180 cgcttctctg gctctggctc gggcacggac tttaccctta ccatcagctc tcttcagccg 240
găagactttğ ccacctatta tiğtcağcag tctaatagct ggccgtatac cttcggtcaa 300
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ggtaccaagg tcgagattaa gcgc
<210> 5
<211> 116
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<223> Description of Artificial Sequence: Synthetic VH Domain
      peptide of Hum B
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Asn Ile Lys Asp Thr
20 25 30
Tyr Ile His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45
Ala Arg Ile Asp Pro Ala Asn Asp Asn Thr Ile Tyr Ala Asp Ser Val 50 55 60
Lys Gly Arg Phe Thr Ile Ser Ser Asp Asp Ser Lys Asn Thr Ala Tyr 65 70 75 80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85
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Thr Ala Ser Gly Tyr Trp Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val
100 105 110
Thr Val Ser Ser
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tcttgcgcag caagcggttt caacattaag gacacctaca tccattgggt gaggcaagct 120
                                                                       180
ccgggtaagg gtctggagtg ggtggcacgt atcgacccgg caaacgacaa caccatttac
gcīgācagēg īgaagggēcg ittiactatt tetagegaeg aetetaagaa cacegegtae 240
                                                                       300
cttcagatga actctctgcg tgccgaggac accgccgtct actactgcac ggcctctggc
                                                                       348
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<210> 7
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<223> Description of Artificial Sequence: Synthetic VL Domain
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Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Asn Asn 20 25 30
Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
35 40 45
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Tyr His Ala Ser Gln Ser Ile Ser Gly Val Pro Ser Arg Phe Ser Gly

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Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Asn Ser Trp Pro Tyr
Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg
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ggtaaagctc cgaaacttct tatctatcac gcctctcaga gcattagcgg cgttccgagc
cgcttctctg gctctggctc gggcacggac tttaccctta ccatcagctc tcttcagccg
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gaagactttg ccacctatta tigtcagcag tctaatagct ggccgtatac cttcggtcaa 300
ggtaccaagg tcgagattaa gcgc
<210> 9
<211> 116
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic VH Domain
      peptide of Hum C
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1 10 15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Asn Ile Lys Asp Thr

· 20 25 30
Tyr Ile His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45
Ala Arg Ile Asp Pro Ala Asn Asp Asn Thr Ile Tyr Ala Asp Ser Val
Lys Gly Arg Phe Thr Ile Ser Gly Asp Asp Ser Lys Asn Thr Ala Tyr
65 70 75 80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
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90
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Thr Thr Ser Gly Tyr Trp Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val
100 105 110
Thr Val Ser Ser
        115
<210> 10
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<211> 348

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<212> DNA
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tcttgcgcag caagcggttt caacattaag gacacctaca tccattgggt gaggcaagct 120
ccgggtaagg gtctggagtg ggtggcacgt atcgacccgg caaacgacaa caccatttac 180
gctgacagcg tgaagggccg titiactait tctggcgacg actctaagaa caccgcgtac 240
cttcagatga actctctgcg tgccgaggac accgccgtct actactgcac gacctctggc 300
tactggtttg cctactgggg ccagggcacg cttgtcaccg tctcttct
<210> 11
<211> 108
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<213> Artificial Sequence
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      peptide of Hum C
<400> 11
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Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Asn Asn
Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
35 40 45
Tyr His Ala Ser Gln Ser Ile Ser Gly Val Pro Ser Arg Phe Ser Gly
50 60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
65 70 75 80
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Asn Ser Trp Pro Tyr
Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg
<210> 12
<211> 324
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic VL nucleotide
      sequence of Hum C
<400> 12
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ggtaaagctc cgaaacttct tatctatcac gcctctcaga gcattagcgg cgttccgagc 180
cgcttctctg gctctggctc gggcacggac tttaccctta ccatcagctc tcttcagccg 240
gaagactttg ccacctatta ttgtcagcag tctaatagct ggccgtatac cttcggtcaa 300
ggtaccaagg tcgagattaa gcgc
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<210> 13
<211> 116
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic VH Domain
      peptide of Hum D
<400> 13
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1 10 15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Asn Ile Lys Asp Thr
20 25 30
Tyr Ile His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
Ala Arg Ile Asp Pro Ala Asn Asp Asn Thr Ile Tyr Ala Asp Ser Val 50 55 60
Lys Gly Arg Phe Thr Ile Ser Ser Asp Asp Ser Lys Asn Thr Ala Tyr
65 70 75 80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Thr Thr Ser Gly Tyr Trp Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val 100 \, 105 \, 110 \,
Thr Val Ser Ser
        115
<210> 14
<211> 348
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<223> Description of Artificial Sequence: Synthetic VH nucleotide
      sequence of Hum D
<400> 14
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tettgegeag caageggttt caacattaag gacacetaca tecattgggt gaggeaaget 120
ccgggtaagg gtctggagtg ggtggcacgt atcgacccgg caaacgacaa caccatttac 180
gctgacagcg tgaagggccg ttttactatt tctagcgacg actctaagaa caccgcgtac
cttcagatga actctctgcg tgccgaggac accgccgtct actactgcac gacctctggc
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                                                                       348
<210> 15
<211> 108
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic VL Domain
      peptide of Hum D
<400> 15
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5 10 15

Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Asn Asn 20 25 30

Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile 35 40 45

Tyr His Ala Ser Gln Ser Ile Ser Gly Val Pro Ser Arg Phe Ser Gly 50 60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro 65 70 75 80

Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Asn Ser Trp Pro Tyr 85 90 95

Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg 100 105

<210> 16

<211> 324

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic VL nucleotide sequence of Hum D

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attacttgcc gcgccagcca atctatcagt aataatcttc actggtatca acaaaaaccg 120
ggtaaagctc cgaaacttct tatctatcac gcctctcaga gcattagcgg cgttccgagc 180
cgcttctctg gctctggctc gggcacggac tttaccctta ccatcagctc tcttcagccg 240
gaagactttg ccacctatta ttgtcagcag tctaatagct ggccgtatac cttcggtcaa 300
ggtaccaagg tcgagattaa gcgc
324

<210> 17 <211> 116

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic VH Domain peptide of Hum E

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Asn Ile Lys Asp Thr 20 25 30

Tyr Ile His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val

Ala Arg Ile Asp Pro Ala Asn Asp Asn Thr Ile Tyr Asp Pro Lys Val

Gln Gly Arg Phe Thr Ile Ser Ala Asp Asp Ser Lys Asn Thr Ala Tyr 65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys

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Thr Thr Ser Gly Tyr Trp Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val 100 105 110

Thr Val Ser Ser 115

<210> 18

<211> 348 <212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic VH nucleotide
 sequence of Hum E

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<210> 19

<211> 108

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic VL Domain peptide of Hum E

Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Asn Asn 20 25 30

Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile 35 40 45

Tyr His Ala Ser Gln Ser Ile Ser Gly Val Pro Ser Arg Phe Ser Gly
50 60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro 65 70 75 80

Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Asn Ser Trp Pro Tyr
85
90
95

Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg 100 105

<210> 20

<211> 324

<212> DNA

<213> Artificial Sequence

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ggtaaagctc cgaaacttct tatctatcac gcctctcaga gcattagcgg cgttccgagc 180 cgcttctctg gctctggctc gggcacggac tttaccctta ccatcagctc tcttcagccg 240 gaagactttg ccacctatta ttgtcagcag tctaatagct ggccgtatac cttcggtcaa 300
ggtaccaagg tcgagattaa gcgc
<210> 21
<211> 116
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        peptide of Hum F
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Asn Ile Lys Asp Thr
20 25 30
Tyr Ile His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45
Ala Arg Ile Asp Pro Ala Asn Asp Asn Thr Ile Tyr Ala Asp Ser Val50 \hspace{1.5cm} 55 \hspace{1.5cm} 60
Lys Gly Arg Phe Thr Ile Ser Ala Asp Asp Ser Lys Asn Thr Ala Tyr
65 70 75 80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85
90
95
Thr Thr Ser Gly Tyr Trp Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val
100 105 110
Thr Val Ser Ser
           115
<210> 22
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        sequence of Hum F
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tcttgcgcag caagcggttt caacattaag gacacctaca tccattgggt gaggcaagct 120 ccgggtaagg gtctggagtg ggtggcacgt atcgacccgg caaacgacaa caccatttac 180 gctgacagcg tgaagggccg ttttactatt tctgcggacg actctaagaa caccgcgtac 240 cttcagatga actctctgcg tgccgaggac accgccgtct actactgcac gacctctggc 300
tactggtttg cctactgggg ccagggcacg cttgtcaccg tctcttct
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<210> 23
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Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Asn Asn 20 25 30
Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
35 40 45
Tyr His Ala Ser Gln Ser Ile Ser Gly Val Pro Ser Arg Phe Ser Gly 50 60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
65 70 75 80
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Asn Ser Trp Pro Tyr
Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg
            100 .
<210> 24
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ggtaaagctc cgaaacttct tatctatcac gcctctcaga gcattagcgg cgttccgagc 180
cgcttctctg gctctggctc gggcacggac tttaccctta ccatcagctc tcttcagccg 240
gaagactttg ccacctatta ttgtcagcag tctaatagct ggccgtatac cttcggtcaa 300
                                                                      324
ggtaccaagg tcgagattaa gcgc
<210> 25
<211> 116
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<213> Artificial Sequence
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      peptide of Hum G
<400> 25
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Asn Ile Lys Asp Thr
20 25 30
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Tyr Ile His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
Ala Arg Ile Asp Pro Ala Asn Asp Asn Thr Ile Tyr Ala Asp Ser Val 50 55 60
Lys Gly Arg Phe Thr Ile Ser Ala Asp Asp Ser Lys Asn Thr Ala Tyr
65 70 75 80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr
85 90 95
Thr Thr Ser Gly Tyr Trp Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val
100 105 110
Thr Val Ser Ser
         115
<210> 26
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tcttgcgcag caagcggttt caacattaag gacacctaca tccattgggt gaggcaagct 120
ccgggtaagg gtctggagtg ggtggcacgt atcgacccgg caaacgacaa caccatttac
gctgacagcg tgaagggccg ttttactatt tctgcggacg actctaagaa caccgcgtac 240 cttcagatga actctctgcg tgccgaggac accgccgtct actactgcac gacctctggc 300
tactggtttg cctactgggg ccagggcacg cttgtcaccg tctcttct
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Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Asn Asn 20 25 30
Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
35 40 45
Lys His Ala Ser Gln Ser Ile Ser Gly Val Pro Ser Arg Phe Ser Gly 50 60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro 65 70 75 80
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Asn Ser Trp Pro Tyr
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Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg
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<223> Description of Artificial Sequence: Synthetic VL nucleotide
      sequence of Hum G
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attacttgcc gcgccagcca atctatcagt aataatcttc actggtatca acaaaaaccg
ggtaaagctc cgaaacttct tatcaaacac gcctctcaga gcattagcgg cgttccgagc 180
cgcttctctg gctctggctc gggcacggac tttaccctta ccatcagctc tcttcagccg 240
găagacttig ccacctătta tigicagcag tctaatagct ggccgtatac cttcggtcaa 300
ggtaccaagg tcgagattaa gcgc
<210> 29
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<213> Artificial Sequence
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      peptide of Hum H
<400> 29
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Asn Ile Lys Asp Thr
20 25 30
Tyr Ile His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45
Ala Arg Ile Asp Pro Ala Asn Asp Asn Thr Ile Tyr Asp Pro Lys Val
Gln Gly Arg Phe Thr Ile Ser Ala Asp Asp Ser Lys Asn Thr Ala Tyr
65 70 75 80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Thr Thr Ser Gly Tyr Trp Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val
100 105 110
Thr Val Ser Ser
        115
<210> 30
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sequence of Hum H

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tettgegeag caageggttt caacattaag gacacetaca tecattgggt gaggeaaget 120
ccgggtaagg gtctggagtg ggtggcacgt atcgacccgg caaacgacaa caccatttac
                                                                       180
gatccgaagg tgcagggccg ttttactatt tctgcggacg actctaagaa caccgcgtac 240 cttcagatga actctctgcg tgccgaggac accgccgtct actactgcac gacctctggc 300
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<210> 31
<211> 108
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<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic VL Domain
      peptide of Hum H
<400> 31
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Asn Asn 20 25 30
Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
35 40 45
Lys His Ala Ser Gln Ser Ile Ser Gly Val Pro Ser Arg Phe Ser Gly
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Asn Ser Trp Pro Tyr
Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg
<210> 32
<211> 324
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic VL nucleotide
      sequence of Hum H
<400> 32
gatatccaga tgacccaatc tccgtctagc ctgagcgcca gtgttggtga tcgagttacc 60
attacttgcc gcgccagcca atctatcagt aataatcttc actggtatca acaaaaaccg 120
ggtaaagctc cgaaacttct tatcaaacac gcctctcaga gcattagcgg cgttccgagc 180
cgcttctctg gctctggctc gggcacggac tttaccctta ccatcagctc tcttcagccg 240
gaagactttg ccacctatta ttgtcagcag tctaatagct ggccgtatac cttcggtcaa 300
ggtaccaagg tcgagattaa gcgc
                                                                       324
<210> 33
<211> 116
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic VH Domain
      peptide of Hum I
<400> 33
Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 10 15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Asn Ile Lys Asp Thr
20 25 30
Tyr Ile His Trp Met Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
Ala Arg Ile Asp Pro Ala Asn Asp Asn Thr Ile Tyr Asp Pro Lys Val
Gln Gly Arg Phe Thr Met Ser Ala Asp Thr Ser Lys Asn Thr Ala Tyr
65 70 75 80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95
Thr Thr Ser Gly Tyr Trp Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val
100 105 110
Thr Val Ser Ser
<210> 34
<211> 348
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic VH nucleotide
      sequence of Hum I
<400> 34
gaagttcaac ttgttgagtc tggtggcggt ctggttcagc cgggtggctc tctgcgcctg 60
tcttgcgcag caagcggttt caacattaag gacacctaca tccattggat gaggcaagct 120
ccgggtaagg gtctggagtg ggtggcacgt atcgacccgg caaacgacaa caccatttac 180
gatccgaagg tgcagggccg ttttactatg tctgcggacg actctaagaa caccgcgtac 240
cttcagatga actctctgcg tgccgaggac accgccgtct actactgcac gacctctggc 300
tactggtttg cctactgggg ccagggcacg cttgtcaccg tctcttct
<210> 35
<211> 108
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic VL Domain
      peptide of Hum I
<400> 35
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly 1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15
Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Asn Asn 20 25 30
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<220>

Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile 35 40 45 Lys His Ala Ser Gln Ser Ile Ser Gly Val Pro Ser Arg Phe Ser Gly 50 60 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro 65 70 75 80 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Asn Ser Trp Pro Tyr 85 90 95 Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg <210> 36 <211> 324 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic VL nucleotide sequence of Hum I <400> 36 gatatccaga tgacccaatc tccgtctagc ctgagcgcca gtgttggtga tcgagttacc 60 attacttgcc gcgccagcca atctatcagt aataatcttc actggtatca acaaaaaccg 120 ggtaaagete egaaacitet tateaaacae geeteteaga geattagegg egiteegage 180 cgcttctctg gctctggctc gggcacggac tttaccctta ccatcagctc tcttcagccg 240 gaagactttg ccacctatta ttgtcagcag tctaatagct ggccgtatac cttcggtcaa 300 ggtaccaagg tcgagattaa gcgc <210> 37

<211> 116

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Murine 1A6 VH Domain

Glu Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Lys Pro Gly Ala 1 1 15 Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Thr Tyr Ile His Trp Met Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile 35 40 45 Gly Arg Ile Asp Pro Ala Asn Asp Asn Thr Ile Tyr Asp Pro Lys Val Gln Gly Lys Ala Thr Met Thr Ala Asp Thr Ser Ser Asn Thr Ala Tyr 65 70 75 80 Leu Gln Leu Ser Ser Leu Thr Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95 Thr Thr Ser Gly Tyr Trp Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val

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<210> 38
<211> 108
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Murine 1A6 VL Domain
<400> 38
Asp Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser Val Thr Pro Gly
Asp Ser Val Ser Leu Ser Cys Arg Ala Ser Gln Ser Ile Ser Asn Asn 20 25 30
Leu His Trp Tyr Gln Gln Lys Ser His Glu Ser Pro Arg Leu Leu Ile
35 40 45
Lys His Ala Ser Gln Ser Ile Ser Gly Ile Pro Ser Arg Phe Ser Gly 50 60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Ser Val Glu Thr 65 70 75 80
Glu Asp Phe Gly Met Phe Phe Cys Gln Gln Ser Asn Ser Trp Pro Tyr
Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg
<210> 39
<211> 93
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Human VH Domain
      consensus sequence of Heavy Chain Subgroup III (Humiii)
<400> 39
Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Asn Phe Ser Trp Val
20 25 30
Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val Ala Ala Asp Ser Val
45
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Asn Thr Ala Tyr 50 60
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
65 70 . 75 80
Thr Arg Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser 85 90
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Thr Val Ser Ala

<210> 40 <211> 81 <212> PRT <213> Artificial Sequence <220>

consensus sequence of Light Chain K Subgroup I (HumKI)

<223> Description of Artificial Sequence: Human VL Domain

Pro Lys Leu Leu Ile Tyr Gly Val Pro Ser Arg Phe Ser Gly Ser Gly 35 40 45

Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro Glu Asp 50 55 60

Phe Ala Thr Tyr Tyr Cys Phe Gly Gln Gly Thr Lys Val Glu Ile Lys 65 70 75 80

Arg

<210> 41 <211> 116 <212> PRT <213> Artificial Sequence

<220> <223> Description of Artificial Sequence: Murine 1A6 VH Domain

\$\frac{400}{61u} \quad \text{Val} & \text{Gln} & \text{Leu} & \text{Gln} & \text{Gln} & \text{Ser} & \text{Gly} & \text{Ala} & \text{Gly} & \text{Pro} & \text{Gly} & \text{Pro} & \text{Gly} & \text{Ala} \\
\text{Ser} & \text{Val} & \text{Lys} & \text{Leu} & \text{Ser} & \text{Cys} & \text{Thr} & \text{Ala} & \text{Ser} & \text{Gly} & \text{Pro} & \text{Glu} & \text{Gln} & \text{Asp} & \text{Pro} & \text{Lys} & \text{Gln} & \text{Asp} & \text{Ala} & \text{Asp} & \text{Thr} & \text{Ala} & \text{Asp} & \text{Thr} & \text{Ala} & \text{Asp} & \text{Thr} & \text{Ser} & \text{Ser} & \text{Asn} & \text{Thr} & \text{Ala} & \text{Val} & \text{Gln} & \text{Asp} & \text{Thr} & \text{Ala} & \text{Val} & \text{Thr} & \text{Ser} & \text{Ser} & \text{Asn} & \text{Thr} & \text{Ala} & \text{Val} & \text{Tyr} & \text{Tyr} & \text{Cys} & \text{Gly} & \text{Gln} & \text{Cys} & \text{Gly} & \text{Thr} & \text{Ala} & \text{Val} & \text{Tyr} & \text{Tyr} & \text{Gly} & \text{Gln} & \text{Gly} & \text{Thr} & \text{Ala} & \text{Val} & \text{Thr} & \text{Val} & \text{Gly} & \text{Gly} & \text{Gln} & \text{Gly} & \text{Gln} & \text{Gly} & \text{Gln} & \text{Gly} & \text{Gln} & \text{Gly} & \text{Gly} & \text{Gly} & \text{Cys} & \text{Gly} & \text{Cys} & \text{Gly} & \text{G

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<211> 108
<212> PRT
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<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Murine 1A6 VL Domain

<400> 42

Asp Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser Val Thr Pro Gly
1 5 10 15

Asp Ser Val Ser Leu Ser Cys Arg Ala Ser Gln Ser Ile Ser Asn Asn 20 25 30

Leu His Trp Tyr Gln Gln Lys Ser His Glu Ser Pro Arg Leu Leu Ile 35 40 45

Lys His Ala Ser Gln Ser Ile Ser Gly Ile Pro Ser Arg Phe Ser Gly 50 60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Ser Val Glu Thr 65 70 75 80

Glu Asp Phe Gly Met Phe Phe Cys Gln Gln Ser Asn Ser Trp Pro Tyr 85 90 95

Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg 100 105

<210> 43

<211> 116 <212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Humanized 1A6 (HumB) VH Domain consensus sequence of Heavy Chain Subgroup III (Humiii)

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Asn Ile Lys Asp Thr 20 25 30

Tyr Ile His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val

Ala Arg Ile Asp Pro Ala Asn Asp Asn Thr Ile Tyr Ala Asp Ser Val 50 55 60

Lys Gly Arg Phe Thr Ile Ser Ser Asp Asp Ser Lys Asn Thr Ala Tyr 65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Thr Ala Ser Gly Tyr Trp Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val 100 105 110

Thr Val Ser Ser 115

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<210> 44
<211> 108
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Humanized 1A6
      (HumB) VL Domain consensus sequence of Light Chain K
      Subgroup I (HumKI)
<400> 44
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Asn Asn 20 25 30
Leu His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
35 40 45
Tyr His Ala Ser Gln Ser Ile Ser Gly Val Pro Ser Arg Phe Ser Gly 50 60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
65 70 75 80
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Asn Ser Trp Pro Tyr
Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg
<210> 45
<211> 93
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Human VH Domain
      consensus sequence of Heavy Chain Subgroup III (Humiii)
<400> 45
Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 10 15
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Asn Phe Ser Trp Val
20 25 30
Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val Ala Ala Asp Ser Val
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Asn Thr Ala Tyr
50 55 60
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
65 70 75 80
Thr Arg Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
```

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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Human VL Domain
      consensus sequence of Light Chain K Subgroup I (HumKI)
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
Asp Arg Val Thr Ile Thr Cys Trp Tyr Gln Gln Lys Pro Gly Lys Ala
Pro Lys Leu Ile Tyr Gly Val Pro Ser Arg Phe Ser Gly Ser Gly 35 40
Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro Glu Asp
Phe Ala Thr Tyr Tyr Cys Phe Gly Gln Gly Thr Lys Val Glu Ile Lys 65 70 75 80
Arg
<210> 47
<211> 753
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic nucleotide
      sequence of Humanized scFv3 (HumI)
<400> 47
cgaaccatgg gcgatatcca gatgacccaa tctccgtcta gcctgagcgc cagtgttggt 60
gatcgagtta ccattacttg ccgcgccagc caatctatca gtaataatct tcactggtat 120
caacaaaaac cgggtaaagc tccgaaactt cttatcaaac acgcctctca gagcattagc 180
ggcgttccga gccgcttctc tggctctggc tcgggcacgg actttaccct taccatcagc 240 tctcttcagc cggaagactt tgccacctat tattgtcagc agtctaatag ctggccgtat 300 accttcggtc aaggtaccaa ggtcgagatt aagcgcggcg gtggcggttc tggtggcggt 360
ggtagcggtg gcggtggatc cggtggcggt ggcagcgaag ttcaacttgt tgagtctggt 420
ggcggtctgg ttcagccggg tggctctctg cgcctgtctt gcgcagcaag cggtttcaac 480
attaaggaca cctacatcca ttggatgagg caagctccgg gtaagggtct ggagtgggtg 540
gcacgtatcg acceggcaaa cgacaacacc atttacgatc cgaaggtgca gggccgtttt 600
actatgtctg cggacacctc taagaacacc gcgtaccttc agatgaactc tctgcgtgcc 660
gaggačaccý cčýtctacta ctyčacyacc tctgyctact gytttyccta ctygygccay
ggcacgcttg tcaccgtctc ttctggttaa ccc
<210> 48
<211> 61
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      AVL-1
<400> 48
cgaaccatgg gcgatatcca gatgacccaa tctccgtcta gcctgagcgc cagtgttggt 60
```

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<210> 49
<211> 72
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     AVL-2
<400> 49
gtgaagatta ttactgatag attggctggc gcggcaagta atggtaactc gatcaccaac 60
actggcgctc ag
<210> 50
<211> 71
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     AVL-3
<400> 50
ctatcagtaa taatcttcac tggtatcaac aaaaaccggg taaagctccg aaacttctta 60
tctatcacgc c
<210> 51
<211> 68
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     AVL-4
<400> 51
cccqaqccaq aqccaqaqaa gcgqctcgga acgccgctaa tgctctgaga ggcgtgatag 60
ataagaag
<210> 52
<211> 70
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      AVL-5
<400> 52
ctctggctct ggctcgggca cggactttac ccttaccatc agctctcttc agccggaaga 60
ctttgccacc
<210> 53
<211> 66
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      AVL-6
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<400> 53
ccttgaccga aggtatacgg ccagctatta gactgctgac aataataggt ggcaaagtct 60
tccggc
<210> 54
<211> 71
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
<400> 54
gtataccttc ggtcaaggta ccaaggtcga gattaagcgc ggcggtggcg gttctggtgg 60
cggtggtagc g
<210> 55
<211> 32
<2:12> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     AVL-8
<400> 55
                                                                    32
cgaaccatgg gcgatatcca gatgacccaa tc
<210> 56
<211> 33
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     AVL-9
<400> 56
                                                                    33
cggatccacc gccaccgcta ccaccgccac cag
<210> 57
<211> 73
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     AVH-1
<400> 57
ggtggcggtg gatccggtgg cggtggcagc gaagttcaac ttgttgagtc tggtggcggt 60
ctggttcagc cgg
<210> 58
<211> 71
<212> DNA
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     AVH-2
<400> 58
gtccttaatg ttgaaaccgc ttgctgcgca agacaggcgc agagagccac ccggctgaac 60
<210> 59
<211> 67
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      AVH-3
<400> 59
ggtttcaaca ttaaggacac ctacatccat tgggtgaggc aagctccggg taagggtctg 60
gagtggg
<210> 60
<211> 76
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     AVH-4
<400> 60
ggcccttcac gctgtcagcg taaatggtgt tgtcgtttgc cgggtcgata cgtgccaccc 60
actccagacc cttacc
<210> 61
<211> 81
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      AVH-5
<400> 61
cgctgacagc gtgaagggcc gttttactat ttctagcgac gactctaaga acaccgcgta 60
ccttcagatg aactctctgc g
<210> 62
<211> 67
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      AVH-6
ccaqtaqcca qaqtccqtqc aqtaqtagac ggcggtgtcc tcggcacgca gagagttcat 60
ctgaagg
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<210> 63
<211> 65
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      AVH-7
<400> 63
ggactctggc tactggtttg cctactgggg ccagggcacg cttgtcaccg tctcttctgg 60
ttaac
<210> 64
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
<400> 64
                                                                     18
ggtggcggtg gatccggt
<210> 65
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      AVH-9
<400> 65
                                                                     20
gggttaacca gaagagacgg
<210> 66
<211> 67
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      BVH-6
<400> 66
ccagtagcca gaggccgtgc agtagtagac ggcggtgtcc tcggcacgca gagagttcat 60
ctgaagg
<210> 67
<211> 65
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      BVH-7
<400> 67
ggcctctggc tactggtttg cctactgggg ccagggcacg cttgtcaccg tctcttctgg 60
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ttaac 65

<210> 68 <211> 81 <212> DNA <213> Artificial Sequence
<220> <223> Description of Artificial Sequence: Synthetic oligonucleotide CVH-5
<400> 68 cgctgacagc gtgaagggcc gttttactat ttctggcgac gactctaaga acaccgcgta 60 ccttcagatg aactctctgc g
<210> 69 <211> 67 <212> DNA <213> Artificial Sequence
<220> <223> Description of Artificial Sequence: Synthetic oligonucleotide CVH-6
<400> 69 ccagtagcca gaggtcgtgc agtagtagac ggcggtgtcc tcggcacgca gagagttcat 60 ctgaagg
<210> 70 <211> 65 <212> DNA <213> Artificial Sequence
<220> <223> Description of Artificial Sequence: Synthetic oligonucleotide CVH-7
<400> 70 gacctctggc tactggtttg cctactgggg ccagggcacg cttgtcaccg tctcttctgg 60 ttaac 65
<210> 71 <211> 67 <212> DNA <213> Artificial Sequence
<220> <223> Description of Artificial Sequence: Synthetic oligonucleotide DVH-6
<400> 71 ccagtagcca gaggtcgtgc agtagtagac ggcggtgtcc tcggcacgca gagagttcat 60 ctgaagg
<210> 72 <211> 65 <212> DNA <213> Artificial Sequence
<220>

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<223> Description of Artificial Sequence: Synthetic oligonucleotide
     DVH-7
<400> 72
gacctctggc tactggtttg cctactgggg ccagggcacg cttgtcaccg tctcttctgg 60
<210> 73
<211> 76
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      EVH-4
<400> 73
ggccctgcac cttcggatcg taaatggtgt tgtcgtttgc cgggtcgata cgtgccaccc 60
actccagacc cttacc
<210> 74
<211> 81
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      EVH-5
<400> 74
cgatccgaag gtgcagggcc gttttactat ttctgcggac gactctaaga acaccgcgta 60
                                                                    81
ccttcagatg aactctctgc g
<210> 75
<211> 67
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      EVH-6
<400> 75
ccaqtaqcca qaqqtcqtqc aqtaqtaqac qqcqqtqtcc tcggcacgca gagagttcat 60
ctgaagg
<210> 76
<211> 65
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      EVH-7
<400> 76
gacctctggc tactggtttg cctactgggg ccagggcacg cttgtcaccg tctcttctgg 60
ttaac
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<210> 77

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<211> 67
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
<400> 77
ccagtagcca gaggtcgtgc agtagtagac ggcggtgtcc tcggcacgca gagagttcat 60
ctgaagg
<210> 78
<211> 65
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      FVH-7
<400> 78
gacctctggc tactggtttg cctactgggg ccagggcacg cttgtcaccg tctcttctgg 60
ttaac
<210> 79
<211> 71
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      GVL-3
<400> 79
ctatcagtaa taatcttcac tggtatcaac aaaaaccggg taaagctccg aaacttctta 60
tcaaacacgc c
<210> 80
<211> 68
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      GVL-4
<400> 80
cccgagccag agccagagaa gcggctcgga acgccgctaa tgctctgaga ggcgtgaaag 60
ataagaag
<210> 81
<211> 81
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      GVH-5
<400> 81
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cgctgacagc gtgaagggcc gttttactat ttctgcggac gactctaaga acaccgcgta 60
ccttcagatg aactctctgc g
<210> 82
<211> .67
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic oligonucleotide
<400> 82
ccaqtaqcca qaqqtcqtqc aqtaqtaqac qqcqqtqtcc tcgqcacgca gagagttcat 60
ctgaagg
<210> 83
<211> 65
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     GVH-7
<400> 83
gacctctggc tactggtttg cctactgggg ccagggcacg cttgtcaccg tctcttctgg 60
<210> 84
<211> 71
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     HVL-3
<400> 84
ctatcagtaa taatcttcac tggtatcaac aaaaaccggg taaagctccg aaacttctta 60
tcaaacacgc c
<210> 85
<211> 68
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     HVL-4
<400> 85
cccqaqccaq aqccaqaqaa qcqqctcqqa acgccqctaa tgctctqaqa ggcgtqaaag 60
ataagaag
<210> 86
<211> 76
<212> DNA
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     HVH-4
<400> 86
ggccctgcac cttcggatcg taaatggtgt tgtcgtttgc cgggtcgata cgtgccaccc 60
actccagacc cttacc
<210> 87
<211> 81
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
     HVH-5
<400> 87
cgatccgaag gtgcagggcc gttttactat ttctgcggac gactctaaga acaccgcgta 60
ccttcagatg aactctctgc g
<210> 88
<211> 67
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      HVH-6
<400> 88
ccagtagcca gaggtcgtgc agtagtagac ggcggtgtcc tcggcacgca gagagttcat 60
ctgaagg
<210> 89
<211> 65
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      HVH-7
<400> 89
qacctctqqc tactqqtttq cctactqqqq ccagggcacg cttgtcaccg tctcttctgg 60
<210> 90
<211> 71
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide
      IVL-3
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